

Exhibit F

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<!--StartFragment-->RESULT 1
US-09-997-722-215
; Sequence 215, Application US/09997722
; Publication No. US20040072154A1
; GENERAL INFORMATION:
; APPLICANT: Morris, David
; APPLICANT: Engelhard, Eric
; TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR CANCER
; FILE REFERENCE: A-71171/RMS/DCF
; CURRENT APPLICATION NUMBER: US/09/997,722
; CURRENT FILING DATE: 2001-11-30
; PRIOR APPLICATION NUMBER: US 09/747,377
; PRIOR FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: US 09/798,586
; PRIOR FILING DATE: 2001-03-02
; NUMBER OF SEQ ID NOS: 301
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 215
; LENGTH: 2119
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-997-722-215

Query Match          100.0%; Score 2119; DB 3; Length 2119;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 2119; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1 GGCACGAGTAGGGGTGGCGGGTCAGTGTGCTCGGGGGCTTCTCCATCCAGGTCCCTGGA 60
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1 GGCACGAGTAGGGGTGGCGGGTCAGTGTGCTCGGGGGCTTCTCCATCCAGGTCCCTGGA 60

Qy     61 GTTCCTGGTCCCTGGAGCTCCGCACTTGGCGCGCAACCTGCGTGAGGCAGCGCGACTCTG 120
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     61 GTTCCTGGTCCCTGGAGCTCCGCACTTGGCGCGCAACCTGCGTGAGGCAGCGCGACTCTG 120

Qy    121 GCGACTGGCCGGCCATGCGCTTCCCGGGCTGAGGACTATGAAGTGGTGTACACCATTTGGCA 180
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    121 GCGACTGGCCGGCCATGCGCTTCCCGGGCTGAGGACTATGAAGTGGTGTACACCATTTGGCA 180

Qy    181 CAGGCTCCTACGGCCGCTGCCAGAAGATCCGGAGGAAGAGTGATGGCAAGATATTAGTTT 240
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    181 CAGGCTCCTACGGCCGCTGCCAGAAGATCCGGAGGAAGAGTGATGGCAAGATATTAGTTT 240

Qy    241 GGAAAGAACTTGACTATGGCTCCATGACAGAAGCTGAGAAACAGATGCTTGTCTTGAAG 300
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    241 GGAAAGAACTTGACTATGGCTCCATGACAGAAGCTGAGAAACAGATGCTTGTCTTGAAG 300

Qy    301 TGAATTGTCTTCGTGAAGTGAACATCCAAACATCGTTCTGTTACTATGATCGGATTATTG 360
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    301 TGAATTGTCTTCGTGAAGTGAACATCCAAACATCGTTCTGTTACTATGATCGGATTATTG 360

Qy    361 ACCGGACCAATACAACACTGTACATTGTAATGGAATATGTGAAGGAGGGGATCTGGCTA 420
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    361 ACCGGACCAATACAACACTGTACATTGTAATGGAATATGTGAAGGAGGGGATCTGGCTA 420

Qy    421 GTGTAATTACAAAGGGAACCAAGGAAAGGCAATACTTAGATGAAGAGTTTGTCTTCGAG 480
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    421 GTGTAATTACAAAGGGAACCAAGGAAAGGCAATACTTAGATGAAGAGTTTGTCTTCGAG 480

Qy    481 TGATGACTCAGTTGACTCTGGCCCTGAAGGAATGCCACAGACGAAGTGATGGTGGTCATA 540
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    481 TGATGACTCAGTTGACTCTGGCCCTGAAGGAATGCCACAGACGAAGTGATGGTGGTCATA 540

Qy    541 CCGTATTGCATCGGGATCTTAAACCAGCCAATGTTTTCTGGATGGCAAGCAAAACGTCA 600
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Db    541 CCGTATTGCATCGGGATCTTAAACCAGCCAATGTTTTCTGGATGGCAAGCAAAACGTCA 600

Qy    601 AGCTTGGAGACTTTGGGCTAGCTAGAATATTAACCATGACACGAGTTTGC AAAACAT 660
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    601 AGCTTGGAGACTTTGGGCTAGCTAGAATATTAACCATGACACGAGTTTGC AAAACAT 660

Qy    661 TTGTTGGCACACCTTATTACATGTCTCCTGAACAAATGAATCGCATGTCTACAATGAGA 720
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    661 TTGTTGGCACACCTTATTACATGTCTCCTGAACAAATGAATCGCATGTCTACAATGAGA 720

Qy    721 AATCAGATATCTGGTCATTGGGCTGCTTGCTGTATGAGTTATGTGCATTAATGCCTCCAT 780
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Db	721	 AATCAGATATCTGGTCATTGGGCTGCTTGCTGTATGAGTTATGTGCATTAATGCCTCCAT	780
Qy	781	TTACAGCTTTTAGCCAGAAAGAACTCGCTGGGAAAATCAGAGAAGGCAAATTCAGGCGAA	840
Db	781	TTACAGCTTTTAGCCAGAAAGAACTCGCTGGGAAAATCAGAGAAGGCAAATTCAGGCGAA	840
Qy	841	TTCCATACCGTTACTCTGATGAATTGAATGAAATTATTACGAGGATGTTAAACTTAAAGG	900
Db	841	TTCCATACCGTTACTCTGATGAATTGAATGAAATTATTACGAGGATGTTAAACTTAAAGG	900
Qy	901	ATTACCATCGACCTTCTGTTGAAGAAATCTTGAGAACCCTTTAATAGCAGATTTGGTTG	960
Db	901	ATTACCATCGACCTTCTGTTGAAGAAATCTTGAGAACCCTTTAATAGCAGATTTGGTTG	960
Qy	961	CAGACGAGCAAAGAAGAAATCTTGAGAGAAGAGGGCGACAATTAGGAGAGCCAGAAAAAT	1020
Db	961	CAGACGAGCAAAGAAGAAATCTTGAGAGAAGAGGGCGACAATTAGGAGAGCCAGAAAAAT	1020
Qy	1021	CGCAGGATTCCAGCCCTGTATTGAGTGAGCTGAAACTGAAGGAAATTCAGTTACAGGAGC	1080
Db	1021	CGCAGGATTCCAGCCCTGTATTGAGTGAGCTGAAACTGAAGGAAATTCAGTTACAGGAGC	1080
Qy	1081	GAGAGCGAGCTCTCAAAGCAAGAGAAGAAAGATTGGAGCAGAAAGAACAGGAGCTTTGTG	1140
Db	1081	GAGAGCGAGCTCTCAAAGCAAGAGAAGAAAGATTGGAGCAGAAAGAACAGGAGCTTTGTG	1140
Qy	1141	TTCGTGAGAGACTAGCAGAGGACAACTGGCTAGAGCAGAAAAATCTGTTGAAGAACTACA	1200
Db	1141	TTCGTGAGAGACTAGCAGAGGACAACTGGCTAGAGCAGAAAAATCTGTTGAAGAACTACA	1200
Qy	1201	GCTTGCTAAAGGAACGGAAGTTCCTGTCTCTGGCAAGTAATCCAGAACTTCTTAATCTTC	1260
Db	1201	GCTTGCTAAAGGAACGGAAGTTCCTGTCTCTGGCAAGTAATCCAGAACTTCTTAATCTTC	1260
Qy	1261	CATCCTCAGTAATTAAGAAGAAAGTTTCATTTTCAGTGGGAAAGTAAAGAGAACATCATGA	1320
Db	1261	CATCCTCAGTAATTAAGAAGAAAGTTTCATTTTCAGTGGGAAAGTAAAGAGAACATCATGA	1320
Qy	1321	GGAGTGAGAAATTCTGAGAGTCAGCTCACATCTAAGTCCAAGTGCAAGGACCTGAAGAAAA	1380
Db	1321	GGAGTGAGAAATTCTGAGAGTCAGCTCACATCTAAGTCCAAGTGCAAGGACCTGAAGAAAA	1380
Qy	1381	GGCTTCACGCTGCCCAGCTGCGGGCTCAAGCCCTGTCAGATATTGAGAAAAATTACCAAC	1440
Db	1381	GGCTTCACGCTGCCCAGCTGCGGGCTCAAGCCCTGTCAGATATTGAGAAAAATTACCAAC	1440
Qy	1441	TGAAAAGCAGACAGATCCTGGGCATGCGCTAGCCAGGTAGAGAGACACAGAGCTGTGTAC	1500
Db	1441	TGAAAAGCAGACAGATCCTGGGCATGCGCTAGCCAGGTAGAGAGACACAGAGCTGTGTAC	1500
Qy	1501	AGGATGTAATATTACCAACCTTTAAAGACTGATATTCAAATGCTGTAGTGTGAATACTT	1560
Db	1501	AGGATGTAATATTACCAACCTTTAAAGACTGATATTCAAATGCTGTAGTGTGAATACTT	1560
Qy	1561	GGCCCCATGAGCCATGCCTTTCTGTATAGTACACATGATATTCGGAATTGGTTTTACTG	1620
Db	1561	GGCCCCATGAGCCATGCCTTTCTGTATAGTACACATGATATTCGGAATTGGTTTTACTG	1620
Qy	1621	TTCTTCAGCAACTATTGTACAAAATGTTACATTTAATTTTCTTTCTTTTAAAGAAC	1680
Db	1621	TTCTTCAGCAACTATTGTACAAAATGTTACATTTAATTTTCTTTCTTTTAAAGAAC	1680
Qy	1681	ATATTATAAAAAGAATACTTTCTTGTTGGGCTTTTAAATCCTGTGTGTGATTACTAGTAG	1740
Db	1681	ATATTATAAAAAGAATACTTTCTTGTTGGGCTTTTAAATCCTGTGTGTGATTACTAGTAG	1740
Qy	1741	GAACATGAGATGTGACATTCTAAATCTTGGGAGAAAAAATAATATTAGGAAAAAATATT	1800
Db	1741	GAACATGAGATGTGACATTCTAAATCTTGGGAGAAAAAATAATATTAGGAAAAAATATT	1800
Qy	1801	TATGCAGGAAGAGTAGCACTCACTGAATAGTTTTAAATGACTGAGTGGTATGCTTACAAT	1860
Db	1801	TATGCAGGAAGAGTAGCACTCACTGAATAGTTTTAAATGACTGAGTGGTATGCTTACAAT	1860
Qy	1861	TGTCATGTCTAGATTTAAATTTTAAAGTCTGAGATTTTAAATGTTTTGAGCTTAGAAAAC	1920

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      |||
Db      1861  TGT CATGCTAGATTAAATTTTAAGTCTGAGATTTTAAATGTTTTTGAGCTTAGAAAAC 1920
      |||
Qy      1921  CCAGTTAGATGCAATTTGGTCATTAATACCATGACATCTTGCTTATAAAATATTCCATTGC 1980
      |||
Db      1921  CCAGTTAGATGCAATTTGGTCATTAATACCATGACATCTTGCTTATAAAATATTCCATTGC 1980
      |||
Qy      1981  TCTGTAGTTCAAATCTGTTAGCTTTGTGAAAATTCATCACTGTGATGTTTGTATTCTTTT 2040
      |||
Db      1981  TCTGTAGTTCAAATCTGTTAGCTTTGTGAAAATTCATCACTGTGATGTTTGTATTCTTTT 2040
      |||
Qy      2041  TTTTTTCTGTTTAAACAGAAATATGAGCTGTCTGTCATTACCTACTTCTTTCCCACTAAA 2100
      |||
Db      2041  TTTTTTCTGTTTAAACAGAAATATGAGCTGTCTGTCATTACCTACTTCTTTCCCACTAAA 2100
      |||
Qy      2101  TAAAAGAATTCTTCAGTTA 2119
      |||
Db      2101  TAAAAGAATTCTTCAGTTA 2119
<!--EndFragment-->
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